

2

OIPE

RAW SEQUENCE LISTING

DATE: 01/15/2002

PATENT APPLICATION: US/10/017,724

TIME: 18:08:11

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\01152002\J017724.raw

P.5

3 <110> APPLICANT: McCarthy, Jeanette
5 <120> TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF VASCULAR DISEASE
7 <130> FILE REFERENCE: MMI-004
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/017,724
C--> 9 <141> CURRENT FILING DATE: 2001-12-14
9 <150> PRIOR APPLICATION NUMBER: US 60/317,178
10 <151> PRIOR FILING DATE: 2001-09-05
12 <150> PRIOR APPLICATION NUMBER: US 60/329,958
13 <151> PRIOR FILING DATE: 2001-10-16
15 <160> NUMBER OF SEQ ID NOS: 11
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 5784
21 <212> TYPE: DNA
22 <213> ORGANISM: Homo Sapiens
24 <400> SEQUENCE: 1
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27 ctgcagggcc ggtctctcgc tccagcagag cctgcgcctt tctgactcgg tccggaacac 180
28 tgaaccagt catcactgca tctttttggc aaaccaggag ctgagctgca ggaggcagga 240
29 tggctctggag gctggtcctg ctggtctgtt ggtgtggcc cagcacgcaa gctggtcacc 300
30 aggacaaaga cagcaccctc gaccttttca gtatcagcaa catcaaccgc aagaccattg 360
31 gcgccaagca gttccgcggg cccgaccccg gcgtgccggc ttaccgcttc gtgcgctttg 420
32 actacatccc accggtgaac gcagatgacc tcagcaagat caccaagatc atgcggcaga 480
33 aggagggtct ctctctcagc gccagctca agcaggacgg caagtccagg ggcagctgt 540
34 tggctctgga gggccccggt ctctcccaga ggcagttcga gatcgtctcc aacggccccg 600
35 cggacacgct ggatctcacc tactggattg acggcaccgg gcatgtggtc tccctggagg 660
36 acgtcggcct ggctgactcg cagtggaaag acgtcaccgt gcaggtggct ggcgagacct 720
37 acagcttgca cgtgggtcgc gacctcatg gaccagttgc tctggacgag cccttctacg 780
38 agcacctgca ggcggaaaag agccgatgt acgtggccaa aggcctctgc agagagagtc 840
39 acttcagggg tttgcttcag aacgtccacc tagtgttga aaactctgtg gaagatattc 900
40 taagcaagaa gggttgccag caaggccagg gagctgagat caacgccatc agtgagaaca 960
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42 ggaggcccga ggtgtgcgaa cgctcgtgcg aggagctggg aaacatggtc caggagctct 1080
43 cggggctcca cgtcctcgtg aaccagctca gcgagaacct caagagagtg tcgaatgata 1140
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54 ccccggtggtc ggctgcact gtcacctgtg ccggtgggat ccgggagcgc acccggtct 1800
55 gcaacagccc tgagcctcag tacggagggg aggcctgctg ggggatgtg caggagcgtc 1860

ENTERED

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Input Set : A:\Seqlist.txt

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56 agatgtgcaa caagaggagc tgccccgtgg atggctgttt atccaacccc tgcttcccgg 1920
57 gagccagtg cagcagcttc cccgatgggt cctggtcag cggtctctgc cctgtgggct 1980
58 tcttgggcaa tggcaccac tgtgaggacc tggacgagt tgccctggtc cccgacatct 2040
59 gcttctccac cagcaagggt cctcgtctgt tcaacactca gcctggcttc cactgcctgc 2100
60 cctgccccgc ccgatacaga gggaaaccag ccgtcggggg cgccctggaa gcagccaaga 2160
61 cggaaaagca agtgtgtgag cccgaaaacc catgcaagga caagacacac aactgccaca 2220
62 agcacgcgga gtgcatctac ctgggtcact tcagcgaccc catgtacaag tgcgagtgcc 2280
63 agacaggcta cgcgggcgac gggctcatct gcggggagga ctcgacctg gacggctggc 2340
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65 cccatctgcc aaattctggg caggaagact ttgacaagga cgggattggc gatgcctgtg 2460
66 atgatgacga tgacaatgac ggtgtgaccg atgagaagga caactgccag ctccctctca 2520
67 atccccgcca ggctgactat gacaaggatg aggttgggga ccgctgtgac aactgccctt 2580
68 acgtgcacaa ccctgcccag atcgacacag acaacaatgg agagggtgac gcctgctccg 2640
69 tggacattga tggggacgat gtcttcaatg aacgagacaa ttgtccctac gtctacaaca 2700
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73 acgccaacca ggctgaccat gacagagacg gccagggcga cgctgtgac cctgatgatg 2940
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78 ttccgcatca aggcaaggag ctggttcaga cagccaactc ggaccccggc atcgtgtgag 3240
79 gttttgacga gtttgggtct gtggacttca gtggcacatt ctacgtaaac actgaccggg 3300
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96 acaagggtca attcaattt gaagtaatgt tttagtaagg agagattaga agacaacagg 4320
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99 catcaaagaa caacatcct tgcaaatggg tgtgacgcgg ttccagatgt ggatttggca 4500
100 aaacctcatt taagtaaaag gttagcagag caaagtgcgg tgctttagct gctgcttgtg 4560
101 ccgttgtggc gtcggggagg ctctgcctg agcttcttc cccagctttg ctgcctgaga 4620
102 ggaaccagag cagacgcaca ggccggaaaa ggcgcatcta acgcgtatct aggttttggg 4680
103 aactgcggac aagttgcttt tacctgattt gatgatacat ttcattaagg ttccagttat 4740
104 aaatatattg ttaatatatta ttaagtgact atagaatgca actccattta ccagtaactt 4800

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105 attttaaata tgcctagtaa cacatatgta gtataatttc tagaaacaaa catctaataa 4860
106 gtatataatc ctgtgaaaat atgaggcttg ataataattag gttgtcacga tgaagcatgc 4920
107 tagaagctgt aacagaatac atagagaata atgaggagtt tatgatggaa ccttaatata 4980
108 taatgttgcc agcgatttta gttcaatatt tgttactggt atctatctgc tgtatatgga 5040
109 attcttttaa ttcaaagcgt gaaaacgaat cagcatttag tcttgccagg cacaccaaat 5100
110 aatcagtcac gtgtaatatg cacaagtttg tttttgtttt tgtttttttt gttggttggt 5160
111 ttttttgctt taagttgcat gatctttctg caggaaatag tcactcatcc cactccacat 5220
112 aaggggttta gtaagagaag tctgtctgtc tgaatgatga tagggggcaa atctttttcc 5280
113 cctttctgtt aatagtcac acatttctat gccaaacagg aacgatccat aacttttagtc 5340
114 ttaatgtaca cattgcattt tgataaaaatt aattttgttg tttcctttga ggttgatcgt 5400
115 tgtgttggtt tgcgtcactt tttacttttt tgcgtgtgga gctgtattcc cgagacaacg 5460
116 aagcgttggg atacttcatt aaatgtagcg actgtcaaca gcgtgcaggt tttctgtttc 5520
117 tgtgttggtg ggtcaaccgt acaatggtgt gggaatgacg atgatgtgaa tatttagaat 5580
118 gtaccatatt tttgtaaat tatttatggt tttctaaaca aatttatcgt ataggttgat 5640
119 gaaacgcat gtgttttgcc aaagactgta aatatttatt tatgtgttca catgggcaaa 5700
120 atttcaccac tgaaaccctg cacttagcta gaacctcatt tttaaagatt aacaacagga 5760
121 aataaattgt aaaaaagggt ttct 5784
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123 <210> SEQ ID NO: 2

124 <211> LENGTH: 1172

125 <212> TYPE: PRT

126 <213> ORGANISM: Homo Sapiens

128 <400> SEQUENCE: 2

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130 1 5 10 15
131 Gln Ala Gly His Gln Asp Lys Asp Thr Thr Phe Asp Leu Phe Ser Ile
132 20 25 30
133 Ser Asn Ile Asn Arg Lys Thr Ile Gly Ala Lys Gln Phe Arg Gly Pro
134 35 40 45
135 Asp Pro Gly Val Pro Ala Tyr Arg Phe Val Arg Phe Asp Tyr Ile Pro
136 50 55 60
137 Pro Val Asn Ala Asp Asp Leu Ser Lys Ile Thr Lys Ile Met Arg Gln
138 65 70 75 80
139 Lys Glu Gly Phe Phe Leu Thr Ala Gln Leu Lys Gln Asp Gly Lys Ser
140 85 90 95
141 Arg Gly Thr Leu Leu Ala Leu Glu Gly Pro Gly Leu Ser Gln Arg Gln
142 100 105 110
143 Phe Glu Ile Val Ser Asn Gly Pro Ala Asp Thr Leu Asp Leu Thr Tyr
144 115 120 125
145 Trp Ile Asp Gly Thr Arg His Val Val Ser Leu Glu Asp Val Gly Leu
146 130 135 140
147 Ala Asp Ser Gln Trp Lys Asn Val Thr Val Gln Val Ala Gly Glu Thr
148 145 150 155 160
149 Tyr Ser Leu His Val Gly Cys Asp Leu Ile Gly Pro Val Ala Leu Asp
150 165 170 175
151 Glu Pro Phe Tyr Glu His Leu Gln Ala Glu Lys Ser Arg Met Tyr Val
152 180 185 190
153 Ala Lys Gly Ser Ala Arg Glu Ser His Phe Arg Gly Leu Leu Gln Asn
154 195 200 205
155 Val His Leu Val Phe Glu Asn Ser Val Glu Asp Ile Leu Ser Lys Lys
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Output Set: N:\CRF3\01152002\J017724.raw

156	210					215					220					
157	Gly	Cys	Gln	Gln	Gly	Gln	Gly	Ala	Glu	Ile	Asn	Ala	Ile	Ser	Glu	Asn
158	225	230					235					240				
159	Thr	Glu	Thr	Leu	Arg	Leu	Gly	Pro	His	Val	Thr	Thr	Glu	Tyr	Val	Gly
160	245					250					255					
161	Pro	Ser	Ser	Glu	Arg	Arg	Pro	Glu	Val	Cys	Glu	Arg	Ser	Cys	Glu	Glu
162	260					265					270					
163	Leu	Gly	Asn	Met	Val	Gln	Glu	Leu	Ser	Gly	Leu	His	Val	Leu	Val	Asn
164	275					280					285					
165	Gln	Leu	Ser	Glu	Asn	Leu	Lys	Arg	Val	Ser	Asn	Asp	Asn	Gln	Phe	Leu
166	290					295					300					
167	Trp	Glu	Leu	Ile	Gly	Gly	Pro	Pro	Lys	Thr	Arg	Asn	Met	Ser	Ala	Cys
168	305	310					315					320				
169	Trp	Gln	Asp	Gly	Arg	Phe	Phe	Ala	Glu	Asn	Glu	Thr	Trp	Val	Val	Asp
170	325					330					335					
171	Ser	Cys	Thr	Thr	Cys	Thr	Cys	Lys	Lys	Phe	Lys	Thr	Ile	Cys	His	Gln
172	340					345					350					
173	Ile	Thr	Cys	Pro	Pro	Ala	Thr	Cys	Ala	Ser	Pro	Ser	Phe	Val	Glu	Gly
174	355					360					365					
175	Glu	Cys	Cys	Pro	Ser	Cys	Leu	His	Ser	Val	Asp	Gly	Glu	Glu	Gly	Trp
176	370					375					380					
177	Ser	Pro	Trp	Ala	Glu	Trp	Thr	Gln	Cys	Ser	Val	Thr	Cys	Gly	Ser	Gly
178	385	390					395					400				
179	Thr	Gln	Gln	Arg	Gly	Arg	Ser	Cys	Asp	Val	Thr	Ser	Asn	Thr	Cys	Leu
180	405					410					415					
181	Gly	Pro	Ser	Ile	Gln	Thr	Arg	Ala	Cys	Ser	Leu	Ser	Lys	Cys	Asp	Thr
182	420					425					430					
183	Arg	Ile	Arg	Gln	Asp	Gly	Gly	Trp	Ser	His	Trp	Ser	Pro	Trp	Ser	Ser
184	435					440					445					
185	Cys	Ser	Val	Thr	Cys	Gly	Val	Gly	Asn	Ile	Thr	Arg	Ile	Arg	Leu	Cys
186	450					455					460					
187	Asn	Ser	Pro	Val	Pro	Gln	Met	Gly	Gly	Lys	Asn	Cys	Lys	Gly	Ser	Gly
188	465	470					475					480				
189	Arg	Glu	Thr	Lys	Ala	Cys	Gln	Gly	Ala	Pro	Cys	Pro	Ile	Asp	Gly	Arg
190	485					490					495					
191	Trp	Ser	Pro	Trp	Ser	Pro	Trp	Ser	Ala	Cys	Thr	Val	Thr	Cys	Ala	Gly
192	500					505					510					
193	Gly	Ile	Arg	Glu	Arg	Thr	Arg	Val	Cys	Asn	Ser	Pro	Glu	Pro	Gln	Tyr
194	515					520					525					
195	Gly	Gly	Lys	Ala	Cys	Val	Gly	Asp	Val	Gln	Glu	Arg	Gln	Met	Cys	Asn
196	530					535					540					
197	Lys	Arg	Ser	Cys	Pro	Val	Asp	Gly	Cys	Leu	Ser	Asn	Pro	Cys	Phe	Pro
198	545	550					555					560				
199	Gly	Ala	Gln	Cys	Ser	Ser	Phe	Pro	Asp	Gly	Ser	Trp	Ser	Cys	Gly	Phe
200	565					570					575					
201	Cys	Pro	Val	Gly	Phe	Leu	Gly	Asn	Gly	Thr	His	Cys	Glu	Asp	Leu	Asp
202	580					585					590					
203	Glu	Cys	Ala	Leu	Val	Pro	Asp	Ile	Cys	Phe	Ser	Thr	Ser	Lys	Val	Pro
204	595					600					605					

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205 Arg Cys Val Asn Thr Gln Pro Gly Phe His Cys Leu Pro Cys Pro Pro
206      610                      615                      620
207 Arg Tyr Arg Gly Asn Gln Pro Val Gly Val Gly Leu Glu Ala Ala Lys
208 625                      630                      635                      640
209 Thr Glu Lys Gln Val Cys Glu Pro Glu Asn Pro Cys Lys Asp Lys Thr
210                      645                      650                      655
211 His Asn Cys His Lys His Ala Glu Cys Ile Tyr Leu Gly His Phe Ser
212                      660                      665                      670
213 Asp Pro Met Tyr Lys Cys Glu Cys Gln Thr Gly Tyr Ala Gly Asp Gly
214                      675                      680                      685
215 Leu Ile Cys Gly Glu Asp Ser Asp Leu Asp Gly Trp Pro Asn Leu Asn
216      690                      695                      700
217 Leu Val Cys Ala Thr Asn Ala Thr Tyr His Cys Ile Lys Asp Asn Cys
218 705                      710                      715                      720
219 Pro His Leu Pro Asn Ser Gly Gln Glu Asp Phe Asp Lys Asp Gly Ile
220                      725                      730                      735
221 Gly Asp Ala Cys Asp Asp Asp Asp Asp Asn Asp Gly Val Thr Asp Glu
222                      740                      745                      750
223 Lys Asp Asn Cys Gln Leu Leu Phe Asn Pro Arg Gln Ala Asp Tyr Asp
224                      755                      760                      765
225 Lys Asp Glu Val Gly Asp Arg Cys Asp Asn Cys Pro Tyr Val His Asn
226      770                      775                      780
227 Pro Ala Gln Ile Asp Thr Asp Asn Asn Gly Glu Gly Asp Ala Cys Ser
228 785                      790                      795                      800
229 Val Asp Ile Asp Gly Asp Asp Val Phe Asn Glu Arg Asp Asn Cys Pro
230                      805                      810                      815
231 Tyr Val Tyr Asn Thr Asp Gln Arg Asp Thr Asp Gly Asp Gly Val Gly
232                      820                      825                      830
233 Asp His Cys Asp Asn Cys Pro Leu Val His Asn Pro Asp Gln Thr Asp
234                      835                      840                      845
235 Val Asp Asn Asp Leu Val Gly Asp Gln Cys Asp Asn Asn Glu Asp Ile
236      850                      855                      860
237 Asp Asp Asp Gly His Gln Asn Asn Gln Asp Asn Cys Pro Tyr Ile Ser
238 865                      870                      875                      880
239 Asn Ala Asn Gln Ala Asp His Asp Arg Asp Gly Gln Gly Asp Ala Cys
240                      885                      890                      895
241 Asp Pro Asp Asp Asp Asn Asp Gly Val Pro Asp Asp Arg Asp Asn Cys
242                      900                      905                      910
243 Arg Leu Val Phe Asn Pro Asp Gln Glu Asp Leu Asp Gly Asp Gly Arg
244      915                      920                      925
245 Gly Asp Ile Cys Lys Asp Asp Phe Asp Asn Asp Asn Ile Pro Asp Ile
246      930                      935                      940
247 Asp Asp Val Cys Pro Glu Asn Asn Ala Ile Ser Glu Thr Asp Phe Arg
248 945                      950                      955                      960
249 Asn Phe Gln Met Val Pro Leu Asp Pro Lys Gly Thr Thr Gln Ile Asp
250                      965                      970                      975
251 Pro Asn Trp Val Ile Arg His Gln Gly Lys Glu Leu Val Gln Thr Ala
252                      980                      985                      990
253 Asn Ser Asp Pro Gly Ile Ala Val Gly Phe Asp Glu Phe Gly Ser Val

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→ Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/017,724

DATE: 01/15/2002

TIME: 18:08:12

Input Set :: A:\Seqlist.txt

Output Set: N:\CRF3\01152002\J017724.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:317 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:318 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:320 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:352 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

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L:452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:454 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

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L:486 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

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L:795 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

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L:868 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:869 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:877 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:891 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:892 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:893 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/017,724

DATE: 01/15/2002

TIME: 18:08:12

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\01152002\J017724.raw

L:960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:961 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:969 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3